

| | 2/3 | |
|------|---|-------------|
| | #94 L 5'-TAAATCTATAACTACAAAAAACACATA-3'EcoRI | |
| | T -CTTAAATCTATAACTACAAAAAACACATACAGGAATTCCATTCAAGAATAGTTCAAACAA | |
| 3 | -GAATTTAGATATTGATGTTTTTTGTGTATGTCCTTAAGGTAAGTTCTTATCAAGTTTGTT | |
| | GAAGATTACAAACTATCAATTTCATACACAATATAAACGATTAAAAGAATGAGATTTCCT | |
| | CTTCTAATGTTTGATAGTTAAAGTATGTGTTATATTTGCTAATTTTCTTACTCTAAAGGA MetArgPhePro | 4 |
| | TCTATTTTACTGCTGTTTTATTCGCTGCTTCCTCCGCTTTAGCTGCTCCAGTCAACACT | |
| | AGATAAAAATGACGACAAAATAAGCGACGAAGGAGGCGAAATCGACGAGGTCAGTTGTGA SerilePheThrAlaValLeuPheAlaAlaSerSerAlaLeuAlaAlaProValAsnThr | 1086 |
| 1087 | ACCACTGAAGATGAAACGGCTCAAATTCCAGCTGAAGCTGTCATCGGTTACTCTGATTTA | |
| | TGGTGACTTCTACTTTGCCGAGTTTAAGGTCGACTTCGACAGTAGCCAATGAGACTAAAT ThrThrGluAspGluThrAlaGlnIleProAlaGluAlaValIleGlyTyrSerAspLeu | 1146 |
| 1147 | GAAGGTGATTTCGATGTTGCTGTTTTGCCATTTTCCAACTCCACCAATAACGGTTTATTG | • |
| | CTTCCACTAAAGCTACAACGACAAAACGGTAAAAGGTTGAGGTGGTTATTGCCAAATAAC GluGlyAspPheAspValAlaValLeuProPheSerAsnSerThrAsnAsnGlyLeuLeu | 1206 64 |
| 1207 | TTTATCAATACTACTATTGCCTCCATTGCTGCTAAAGAAGAAGGTGTTTCTTTGGATAAA | |
| | AAATAGTTATGATGATAACGGAGGTAACGACGATTTCTTCTTCCACAAAGAAACCTATTT PheIleAsnThrThrIleAlaSerIleAlaAlaLysGluGluGlyValSerLeuAspLys 3'-CCACAAAGAAACCTATTT | 1266 84 |
| 1267 | · · · · · · · · · · · · · · · · · · · | 1326 |
| | TCTAAGCAATTGGTTGTGAACACGCCAAGGGTGAACCAACTTCGAAACATGAACCAAACG ArgPheValAsnGlnHisLeuCysGlySerHisLeuValGluAlaLeuTyrLeuValCys TCT GCAATTGGTTGTGAACACGCCAAGGGTGAACCAACTTCGAAACATGAACC-5' C A #593 1 T ATGTAGCCTTTGGT | 104 |
| | T TGACGATGCT CTTCGACTTCGAC C A | |
| 1327 | GGTGAAAGAGGTTTCTTCTACACTCCTAAG AGGTATTG-3' GGTGAAAGAGGTTTCTTCTACACTCCTAAGGCTGCTAAGGGTATTGTCGAACAATGCTGT | 1206 |
| | CCACTTTCTCCAAAGAAGATGTGAGGATTCCGACGATTCCCATAACAGCTTGTTACGACA GlyGluArgGlyPhePheTyrThrProLysAlaAlaLysGlyIleValGluGlnCysCys | 1386 124 |
| | ACCTCCATCTGCTCCTTGTACCAATTGGAAAACTACTGCAACTAGACGCAGCCCGCAGGC TGGAGGTACACCACCAACAACAACTACTGCAACTAGACGCAGCCCGCAGGC | |
| 7 | TGGAGGTACACCACCAACAACAACAACAAACAAAAAAAAA | 446 |

TGGAGGTAGACGAGGAACATGGTTAACCTTTTGATGACGTTGATCTGCGTCGGGCGTCCG
ThrSerIleCysSerLeuTyrGlnLeuGluAsnTyrCysAsn*** 3'-GGGCGTCCG 138